

Patent Application US/07/996,771

SEQUENCE LISTING

ENTERED

(1) GENERAL INFORMATION:

(i) APPLICANT: R. Bruce Wallace

(ii) TITLE OF INVENTION: Ligation Amplification of
Nucleic Acid Sequences

(iii) NUMBER OF SEQUENCES: 15

(iv) CORRESPONDENCE ADDRESS:

(A) ADDRESSEE: City of Hope

(B) STREET: 1500 East Duarte Road

(C) CITY: Duarte

(D) STATE: California

(E) COUNTRY: United States of America

(F) ZIP: 91010-0269

(v) COMPUTER READABLE FORM:

(A) MEDIUM TYPE: 3M Double Density 5 1/4" diskette

(B) COMPUTER: Wang PC

(C) OPERATING SYSTEM: MS-DOS (R) Version 3.30

(D) SOFTWARE: Microsoft (R)

(vi) CURRENT APPLICATION DATA:

(A) APPLICATION NUMBER: US/07/996,771

(B) FILING DATE: 24 DEC 1992

(C) CLASSIFICATION: 435

(vii) PRIOR APPLICATION DATA:

(A) APPLICATION NUMBER: US/07/870,221

(B) FILING DATE: 20 April 1992

(C) CLASSIFICATION: Unknown

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53
54 (D) APPLICATION NUMBER: US/07/178,377
55
56 (B) FILING DATE: 06 April 1988
57
58 (C) CLASSIFICATION: Unknown
59
60 (viii) ATTORNEY/AGENT INFORMATION:
61
62 (A) NAME: Irons, Edward S.
63
64 (B) REGISTRATION NUMBER: 16,541
65
66 (C) REFERENCE/DOCKET NUMBER: None
67
68 (ix) TELECOMMUNICATION INFORMATION:
69
70 (A) TELEPHONE: (202) 785-6938
71
72 (B) TELEFAX: (202) 785-5351
73
74 (C) TELEX: 440087 LM WSH
75
76
77 (2) INFORMATION FOR SEQ ID NO: 1:
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79 (i) SEQUENCE CHARACTERISTICS:
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81 (A) LENGTH: 14
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83 (B) TYPE: nucleic acid
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85 (C) STRANDEDNESS: Single Stranded
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87 (D) TOPOLOGY: Linear
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89 (ii) MOLECULE TYPE: Not Applicable
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91 (iii) HYPOTHETICAL: Not Applicable
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93 (iv) ANTI-SENSE: Not Applicable
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95 (v) FRAGMENT TYPE: Not Applicable
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97 (vi) ORIGINAL SOURCE: Synthetically Prepared
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99 (vii) IMMEDIATE SOURCE: Synthetically Prepared
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101 (viii) POSITION IN GENOME: None
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103 (ix) FEATURE: None
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105 (x) PUBLICATION INFORMATION: None
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109 CGGCAGACTT CTCC 14
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113 (2) INFORMATION FOR SEQ ID NO: 2:
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115 (i) SEQUENCE CHARACTERISTICS:
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117 (A) LENGTH: 8
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119 (B) TYPE: nucleic acid
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121 (C) STRANDEDNESS: Single Stranded
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123 (D) TOPOLOGY: Linear
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125 (ii) MOLECULE TYPE: Not Applicable
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145 TCAGGAGT 8
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151 (i) SEQUENCE CHARACTERISTICS:
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153 (A) LENGTH: 6
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155 (B) TYPE: nucleic acid
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157 (C) STRANDEDNESS: Single Stranded
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159 (D) TOPOLOGY: Linear
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161 (ii) MOLECULE TYPE: Not Applicable
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163 (iii) HYPOTHETICAL: Not Applicable
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183 (2) INFORMATION FOR SEQ ID NO:4:
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187 (A) LENGTH: 8
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205 (vii) IMMEDIATE SOURCE: Synthetically Prepared
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207 (viii) POSITION IN GENOME: None
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209 (ix) FEATURE: None
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211 (x) PUBLICATION INFORMATION: None
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264 (B) TYPE: nucleic acid
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268 (D) TOPOLOGY: Linear
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270 (ii) MOLECULE TYPE: Not Applicable
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304 (D) TOPOLOGY: Linear
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306 (ii) MOLECULE TYPE: Not Applicable
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308 (iii) HYPOTHETICAL: Not Applicable
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310 (iv) ANTI-SENSE: Not Applicabl
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312 (v) FRAGMENT TYPE: Not Applicable

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313
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342 (ii) MOLECULE TYPE: Not Applicable
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362 GGAGAAGT 8
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384 (v) FRAGMENT TYPE: Not Applicable
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398 GGAGAA 6
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404 (i) SEQUENCE CHARACTERISTICS:
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406 (A) LENGTH: 19
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408 (B) TYPE: nucleic acid
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412 (D) TOPOLOGY: Linear
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414 (ii) MOLECULE TYPE: Not Applicable
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416 (iii) HYPOTHETICAL: Not Applicable

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418 (iv) ANTI-SENSE: Not Applicable
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420 (v) FRAGMENT TYPE: Not Applicable
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422 (vi) ORIGINAL SOURCE: Synthetically Prepared
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426 (viii) POSITION IN GENOME: None
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434 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 10:
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436 CTCCTGAGGA GAAGTCTGC 19
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443 (A) LENGTH: 19
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445 (B) TYPE: nucleic acid
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447 (C) STRANDEDNESS: Single Stranded
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449 (D) TOPOLOGY: Linear
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451 (ii) MOLECULE TYPE: Not Applicable
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457 (v) FRAGMENT TYPE: Not Applicable
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463 (viii) POSITION IN GENOME: None
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465 (ix) FEATURE: None
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467 (x) PUBLICATION INFORMATION: None
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469 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 11:
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471 CTCCTGTGGA GAAGTCTGC 19
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477 (i) SEQUENCE CHARACTERISTICS:
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479 (A) LENGTH: 23
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481 (B) TYPE: nucleic acid
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483 (C) STRANDEDNESS: Single Stranded
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485 (D) TOPOLOGY: Linear
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487 (ii) MOLECULE TYPE: Not Applicable
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489 (iii) HYPOTHETICAL: Not Applicable
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493 (v) FRAGMENT TYPE: Not Applicable
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497 (vii) IMMEDIATE SOURCE: Synthetically Prepared
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499 (viii) POSITION IN GENOME: None
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501 (ix) FEATURE: None
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507 CGGCAGACTT CTCCTCAGGA GTC 23
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515 (A) LENGTH: 23
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517 (B) TYPE: nucleic acid
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519 (C) STRANDEDNESS: Single Stranded
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521 (D) TOPOLOGY: Linear
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523 (ii) MOLECULE TYPE: Not Applicable
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535 (viii) POSITION IN GENOME: None
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539 (x) PUBLICATION INFORMATION: None
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547 (2) INFORMATION FOR SEQ ID NO:14:

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553 (B) TYPE: nucleic acid
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555 (C) STRANDEDNESS: Single Stranded
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571 (viii) POSITION IN GENOME: None
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573 (ix) FEATURE: None
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587 (A) LENGTH: 8
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593 (D) TOPOLOGY: Linear
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609 (ix) FEATURE: None
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611 (x) PUBLICATION INFORMATION: None
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615 TGAGGACT 8
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SEQUENCE VERIFICATION REPORT
PATENT APPLICATION US/07/996,771

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LINE ERROR

ORIGINAL TEXT

51 Unknown or Misplaced Identifier
58 Unknown or Misplaced Identifier

(C) CLASSIFICATION: Unknown
(C) CLASSIFICATION: Unknown

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SEQUENCE MISSING ITEM REPORT
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DATE: 02/02/93
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S381

MANDATORY IDENTIFIER THAT WAS NOT FOUND

PAGE: 1

SEQUENCE CORRECTION REPORT
PATENT APPLICATION US/07/996,771

DATE: 02/02/93
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LINE ORIGINAL TEXT

CORRECTED TEXT